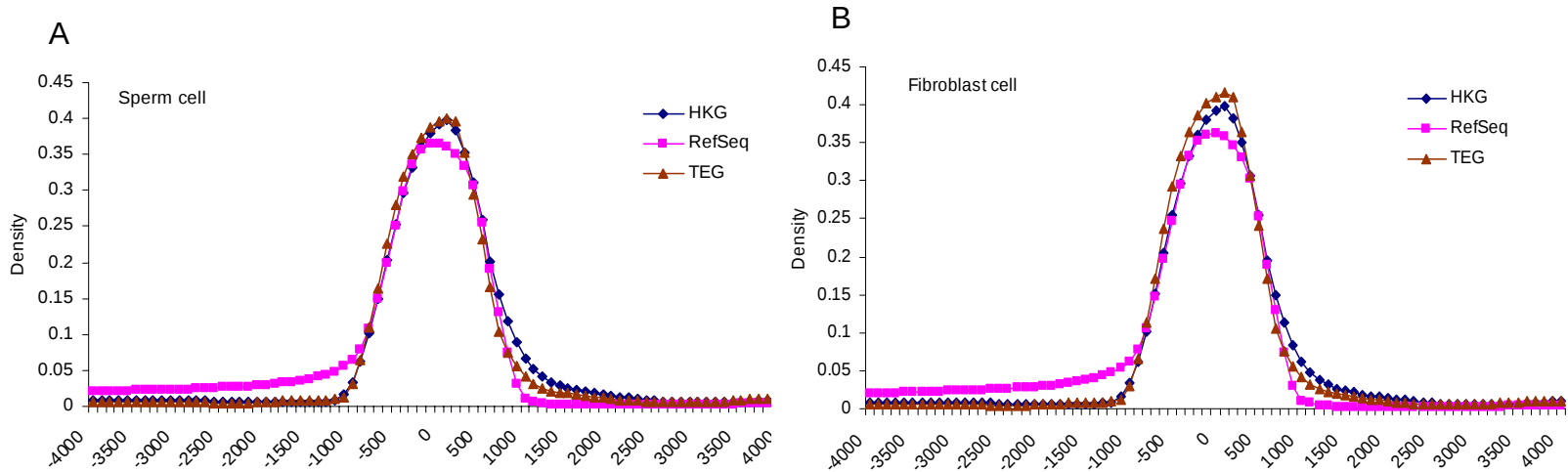




Additional file 6. DNA methylation at transcription start sites.

DNA methylation was detected in sperm cells (A) and fibroblast cells (B) (Weber et al. 2007). The average density of DNA methylation in three gene groups, HKGs, RefSeq genes and TEGs, are calculated in each 500 bp window 4 kb around the transcription start sites.